#### **AMENDMENTS**

# In the Specification

Please amend Table A, page 19, lines 22 through 32 as follows:

### Table A

### **Conservative Substitutions I**

SIDE CHAIN	<b>AMINO ACID</b>		
Aliphatic	Non-polar	G, A, P	
		I, L, V	
	Polar - uncharged	C, S, T, M	
		N, Q	
	Polar - charged	D, E	
		K, R	
Aromatic		H, F, W, Y	
Other		N, Q, D, E	

Please amend Table B, page 21, lines 1 through 19 to read:

**SIDE CHAIN** 

Negatively Charged (Acidic):

## Table B **Conservative Substitutions II**

<b>CHARACTERISTIC</b>	<b>AMINO ACID</b>	
Non-polar (hydrophobic)		
A. Aliphatic:	A, L, I, V, P	
B. Aromatic:	F, W	
C. Sulfur-containing:	M	
D. Borderline:	G	
Uncharged-polar		
A. Hydroxyl:	S, T, Y	
B. Amides:	N, Q	
C. Sulfhydryl:	C	
D. Borderline:	G	
Positively Charged (Basic):	K, R, H	

Please amend Table 1, page 37, line 25 through page 38, line 35 to read as

D, E

follows:



Nucleotide SEQ ID NO:	Representative Isolate	PossibleGene Function	Vaccination # survivors/total	Challenge # survivors/total	LD <sub>50</sub>
	wild type	-	0/10	-	<10
23	PM1B1	guaB	10/10, 10/10, 10/10	9/10, 9/10	4.3 x 10E6
11	PM1D1	dsbB	10/10, 5/10	10/10, 5/5	8.4 x 10E4